



1

SEQUENCE LISTING

MAY 18 2000

TECH CENTER 1600/2900

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Whitney, Michael
Cubitt, Andrew B.
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<120> METHODS OF PROTEIN DESTABILIZATION AND
USES THEREOF

<130> AURO1330

<140> 09/498,098
<141> 2000-02-04

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cagagggttga tctttgtctgg gaaacagctg gaagatggac gcaccctgtc tgactacaac 180
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caa gag caa ctc ggt cgc cgc ata cac tat tct cag aat gac ttg gtt Gln Glu Gln Leu Gly Arg Arg Ile His Tyr Ser Gln Asn Asp Leu Val 65 70 75 80	240
gag tac tca cca gtc aca gaa aag cat ctt acg gat ggc atg aca gta Glu Tyr Ser Pro Val Thr Glu Lys His Leu Thr Asp Gly Met Thr Val 85 90 95	288
aga gaa tta tgc agt gct gcc ata acc atg agt gat aac act gcg gcc Arg Glu Leu Cys Ser Ala Ala Ile Thr Met Ser Asp Asn Thr Ala Ala 100 105 110	336
aac tta ctt ctg aca acg atc gga gga ccg aag gag cta acc gct ttt Asn Leu Leu Thr Thr Ile Gly Gly Pro Lys Glu Leu Thr Ala Phe 115 120 125	384
ttg cac aac atg ggg gat cat gta act cgc ctt gat cgt tgg gaa ccg Leu His Asn Met Gly Asp His Val Thr Arg Leu Asp Arg Trp Glu Pro 130 135 140	432
gag ctg aat gaa gcc ata cca aac gac gag cgt gac acc acg atg cct Glu Leu Asn Glu Ala Ile Pro Asn Asp Glu Arg Asp Thr Thr Met Pro 145 150 155 160	480
gca gca atg gca aca acg ttg cgc aaa cta tta act ggc gaa cta ctt Ala Ala Met Ala Thr Thr Leu Arg Lys Leu Leu Thr Gly Glu Leu Leu 165 170 175	528
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gct gat aaa tct gga gcc ggt gag cgt ggg tct cgc ggt atc att gca Ala Asp Lys Ser Gly Ala Gly Glu Arg Gly Ser Arg Gly Ile Ile Ala 210 215 220	672
gca ctg ggg cca gat ggt aag ccc tcc cgt atc gta gtt atc tac acg Ala Leu Gly Pro Asp Gly Lys Pro Ser Arg Ile Val Val Ile Tyr Thr 225 230 235 240	720
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gat gct gaa gat cag ttg ggt gca cga gtg ggt tac atc gaa ctg gat Asp Ala Glu Asp Gln Leu Gly Ala Arg Val Gly Tyr Ile Glu Leu Asp 35 40 45	144
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gat ggc atg aca gta aga gaa tta tgc agt gct gcc ata acc atg agt Asp Gly Met Thr Val Arg Glu Leu Cys Ser Ala Ala Ile Thr Met Ser 115 120 125	384
gat aac act gcg gcc aac tta ctt ctg aca acg atc gga gga ccg aag Asp Asn Thr Ala Ala Asn Leu Leu Leu Thr Thr Ile Gly Gly Pro Lys 130 135 140	432
gag cta acc gct ttt ttg cac aac atg ggg gat cat gta act cgc ctt Glu Leu Thr Ala Phe Leu His Asn Met Gly Asp His Val Thr Arg Leu 145 150 155 160	480
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165

170

175

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180 185 190

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195 200 205

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210 215 220

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Ala Gly Trp Phe Ile Ala Asp Lys Ser Gly Ala Gly Glu Arg Gly Ser
225 230 235 240

cgc ggt atc att gca gca ctg ggg cca gat ggt aag ccc tcc cgt atc 768
Arg Gly Ile Ile Ala Ala Leu Gly Pro Asp Gly Lys Pro Ser Arg Ile
245 250 255

gta gtt atc tac acg acg ggg agt cag gca act atg gat gaa cga aat 816
Val Val Ile Tyr Thr Gly Ser Gln Ala Thr Met Asp Glu Arg Asn
260 265 270

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ttg ggt gca cga gtg ggt tac atc gaa ctg gat ctc aac agc ggt aag 96
Leu Gly Ala Arg Val Gly Tyr Ile Glu Leu Asp Leu Asn Ser Gly Lys
20 25 30

atc ctt gag agt ttt cgc ccc gaa gaa cgt ttt cca atg atg agc act 144
Ile Leu Glu Ser Phe Arg Pro Glu Glu Arg Phe Pro Met Met Ser Thr
35 40 45

ttt aaa gtt ctg cta tgt ggc gcg gta tta tcc cgt gat gac gcc ggg 192
Phe Lys Val Leu Leu Cys Gly Ala Val Leu Ser Arg Asp Asp Ala Gly
50 55 60

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Gln Glu Gln Leu Gly Arg Arg Ile His Tyr Ser Gln Asn Asp Leu Val				
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Glu Tyr Ser Pro Val Thr Glu Lys His Leu Thr Asp Gly Met Thr Val				
85	90	95		
aga gaa tta tgc agt gct gcc ata acc atg agt gat aac act gcg gcc				336
Arg Glu Leu Cys Ser Ala Ala Ile Thr Met Ser Asp Asn Thr Ala Ala				
100	105	110		
aac tta ctt ctg aca acg atc gga gga ccg aag gag cta acc gct ttt				384
Asn Leu Leu Leu Thr Thr Ile Gly Gly Pro Lys Glu Leu Thr Ala Phe				
115	120	125		
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Leu His Asn Met Gly Asp His Val Thr Arg Leu Asp His Trp Glu Pro				
130	135	140		
gag ctg aat gaa gcc ata cca aac gac gag cgt gac acc acg atg cct				480
Glu Leu Asn Glu Ala Ile Pro Asn Asp Glu Arg Asp Thr Thr Met Pro				
145	150	155	160	
gta gca atg gca aca acg ttg cgc aaa cta tta act ggc gaa cta ctt				528
Val Ala Met Ala Thr Thr Leu Arg Lys Leu Leu Thr Gly Glu Leu Leu				
165	170	175		
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Thr Leu Ala Ser Arg Gln Gln Leu Ile Asp Trp Met Glu Ala Asp Lys				
180	185	190		
gtt gca gga cca ctt ctg cgc tcg gcc ctt ccg gct ggc tgg ttt att				624
Val Ala Gly Pro Leu Leu Arg Ser Ala Leu Pro Ala Gly Trp Phe Ile				
195	200	205		
gct gat aaa tct gga gcc ggt gag cgt ggg tct cgc ggt atc att gca				672
Ala Asp Lys Ser Gly Ala Gly Glu Arg Gly Ser Arg Gly Ile Ile Ala				
210	215	220		
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Ala Leu Gly Pro Asp Gly Lys Pro Ser Arg Ile Val Val Ile Tyr Thr				
225	230	235	240	
acg ggg agt cag gca act atg gat gaa cga aat aga cag atc gct gag				768
Thr Gly Ser Gln Ala Thr Met Asp Glu Arg Asn Arg Gln Ile Ala Glu				
245	250	255		
ata ggt gcc tca ctg att aag cat tgg				795
Ile Gly Ala Ser Leu Ile Lys His Trp				
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 ggt gca cga gtg ggt tac atc gaa ctg gat ctc aac agc ggt aag atc 96
 Gly Ala Arg Val Gly Tyr Ile Glu Leu Asp Leu Asn Ser Gly Lys Ile
 20 25 30

 ctt gag agt ttt cgc ccc gaa gaa cgt ttt cca atg atg agc act ttt 144
 Leu Glu Ser Phe Arg Pro Glu Glu Arg Phe Pro Met Met Ser Thr Phe
 35 40 45

 aaa gtt ctg cta tgt ggc gcg gta tta tcc cgt att gac gcc ggg caa 192
 Lys Val Leu Leu Cys Gly Ala Val Leu Ser Arg Ile Asp Ala Gly Gln
 50 55 60

 gag caa ctc ggt cgc cgc ata cac tat tct cag aat gac ttg gtt gag 240
 Glu Gln Leu Gly Arg Arg Ile His Tyr Ser Gln Asn Asp Leu Val Glu
 65 70 75 80

 tac tca cca gtc aca gaa aag cat ctt acg gat ggc atg aca gta aga 288
 Tyr Ser Pro Val Thr Glu Lys His Leu Thr Asp Gly Met Thr Val Arg
 85 90 95

 gaa tta tgc agt gct gcc ata acc atg agt gat aac act gcg gcc aac 336
 Glu Leu Cys Ser Ala Ala Ile Thr Met Ser Asp Asn Thr Ala Ala Asn
 100 105 110

 tta ctt ctg aca acg atc gga gga ccg aag gag cta acc gct ttt ttg 384
 Leu Leu Leu Thr Thr Ile Gly Gly Pro Lys Glu Leu Thr Ala Phe Leu
 115 120 125

 cac aac atg ggg gat cat gta act cgc ctt gat cat tgg gaa ccg gag 432
 His Asn Met Gly Asp His Val Thr Arg Leu Asp His Trp Glu Pro Glu
 130 135 140

 ctg aat gaa gcc ata cca aac gac gag cgt gac acc acg atg cct gta 480
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 145 150 155 160

 gca atg gca aca acg ttg cgc aaa cta tta act ggc gaa cta ctt act 528
 Ala Met Ala Thr Thr Leu Arg Lys Leu Leu Thr Gly Glu Leu Leu Thr
 165 170 175

 cta gct tcc cgg caa caa tta ata gac tgg atg gag gcg gat aaa gtt 576
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 180 185 190

 gca gga cca ctt ctg cgc tcg gcc ctt ccg gct ggc tgg ttt att gct 624
 Ala Gly Pro Leu Leu Arg Ser Ala Leu Pro Ala Gly Trp Phe Ile Ala
 195 200 205

 gat aaa tct gga gcc ggt gag cgt ggg tct cgc ggt atc att gca gca 672
 Asp Lys Ser Gly Ala Gly Glu Arg Gly Ser Arg Gly Ile Ile Ala Ala
 210 215 220

ctg ggg cca gat ggt aag ccc tcc cgt atc gta gtt atc tac acg acg Leu Gly Pro Asp Gly Lys Pro Ser Arg Ile Val Val Ile Tyr Thr Thr 225 230 235 240	720
ggg agt cag gca act atg gat gaa cga aat aga cag atc gct gag ata Gly Ser Gln Ala Thr Met Asp Glu Arg Asn Arg Gln Ile Ala Glu Ile 245 250 255	768
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ggc gtg ctt ttg caa cag aaa tca ata gaa gat ctg aac cag aga ata Gly Val Leu Leu Gln Gln Lys Ser Ile Glu Asp Leu Asn Gln Arg Ile 50 55 60	192
aca tat aca cgt gat gat ctt gta aac tac aac ccg att acg gaa aag Thr Tyr Thr Arg Asp Asp Leu Val Asn Tyr Asn Pro Ile Thr Glu Lys 65 70 75 80	240
cac gtt gat acg gga atg acg ctc aaa gag ctt gcg gat gct tcg ctt His Val Asp Thr Gly Met Thr Leu Lys Glu Leu Ala Asp Ala Ser Leu 85 90 95	288
cga tat agt gac aat gcg gca cag aat ctc att ctt aaa caa att ggc Arg Tyr Ser Asp Asn Ala Ala Gln Asn Leu Ile Leu Lys Gln Ile Gly 100 105 110	336
gga cct gaa agt ttg aaa aag gaa ctg agg aag att ggt gat gag gtt Gly Pro Glu Ser Leu Lys Lys Glu Leu Arg Lys Ile Gly Asp Glu Val 115 120 125	384
aca aat ccc gaa cga ttc gaa cca gag tta aat gaa gtg aat ccg ggt Thr Asn Pro Glu Arg Phe Glu Pro Glu Leu Asn Glu Val Asn Pro Gly 130 135 140	432

gaa act cag gat acc agt aca gca aga gca ctt gtc aca agc ctt cga Glu Thr Gln Asp Thr Ser Thr Ala Arg Ala Leu Val Thr Ser Leu Arg 145 150 155 160	480
gcc ttt gct ctt gaa gat aaa ctt cca agt gaa aaa cgc gag ctt tta Ala Phe Ala Leu Glu Asp Lys Leu Pro Ser Glu Lys Arg Glu Leu Leu 165 170 175	528
atc gat tgg atg aaa cga aat acc act gga gac gcc tta atc cgt gcc Ile Asp Trp Met Lys Arg Asn Thr Thr Gly Asp Ala Leu Ile Arg Ala 180 185 190	576
gga gcg gca tca tat gga acc cgg aat gac att gcc atc att tgg ccg Gly Ala Ala Ser Tyr Gly Thr Arg Asn Asp Ile Ala Ile Ile Trp Pro 195 200 205	624
cca aaa gga gat cct gtc ggt gtg ccg gac ggt tgg gaa gtg gct gat Pro Lys Gly Asp Pro Val Gly Val Pro Asp Gly Trp Glu Val Ala Asp 210 215 220	672
aaa act gtt ctt gca gta tta tcc agc agg gat aaa aag gac gcc aag Lys Thr Val Leu Ala Val Leu Ser Ser Arg Asp Lys Lys Asp Ala Lys 225 230 235 240	720
tat gat gat aaa ctt att gca gag gca aca aag gtg gta atg aaa gcc Tyr Asp Asp Lys Leu Ile Ala Glu Ala Thr Lys Val Val Met Lys Ala 245 250 255	768
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<212> DNA
<213> Anemonia majano

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<212> DNA

<213> Zoanthus sp.

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ttgtcagctg	cctttaacta	cgaaa	acagg	gttttca	cgaagacata	240
gactattca	agaactcgt	tcctg	ctgg	tatacatgg	agacatagtt	300
gatggagcag	tttgcata	at	aatgc	ataacagt	gtgttgaaga	360
tatcatgagt	ccaaatttta	tgg	atgt	tttccctgctg	atggacctgt	420
atgacagata	actgggagcc	atc	cctgc	aaagatcata	cagtaccta	480
ttgaaagggg	atgtctccat	gtac	ctc	ctgaaggat	gtgggcgtt	540
ttcgacacag	tttaca	aaag	ctgt	ccaagaaaga	tgccggactg	600
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<210> 11

<211> 696

<212> DNA

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gactattca	agaactcgt	tcctg	ctgg	tatacatgg	agacatagtt	300
gatggagcag	tctgcata	caatgt	at	aaacagt	gtgtcaaaga	360
tatcataaga	gcatatttta	tgg	atgt	tttccctgctg	atggacctgt	420
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<211> 699

<212> DNA

<213> Discosoma striata

<400> 12

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accaataccg	tcacgctcga	ggttaccaag	ggtggacctc	tgccatttgg	ttggcatatt	180
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gattatctaa	agctgtcatt	tccggaggga	tatacatggg	aacggtccat	gcactttgaa	300
gacggtggct	tgtgttgat	caccaatgtat	atcagttga	caggcaactg	tttctactac	360
gacatcaagt	tcactggctt	gaactttcct	ccaaatggac	ccgttgtgca	gaagaagaca	420
actggctggg	aaccgagcac	tgagcggtt	tatcctcggt	atgggtgttt	gataggagac	480
atccatcatg	ctctgacagt	tgaaggaggt	ggtcattacg	catgtgacat	taaaactgtt	540
tacagggcca	agaaggccgc	cttgaagatg	ccagggtatac	actatgttga	caccaaactg	600
gttatatgga	acaacgacaa	agaattcatg	aaagttgagg	agcatgaaat	cgccggttgca	660
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His Gly Ser Gly Ala Trp Leu His Pro Glu Thr Leu Val Lys Val Lys
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<212> PRT

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Leu Val Leu Arg Leu Arg Gly Val Gly Ser Val Gly Ala Val Gly Ser
1 5 10 15
Val Gly Asp Glu Val Asp Gly Ser Gly Ala Trp Leu His Pro Glu Thr
20 25 30
Leu Val Lys Val
35